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>SARS120.CB_C03_-_018.ab1 856 40 361 ABT GTACTCGTCATATCGGTGCCTGTGTAGGATGAAGATCGTAATGATCTTAC TACAGTCCTTAATTAGTCCAGTTACATTTTCTGCTTGTAATGTAGCCACA TTGCGACGTGGTATTTCTAGACTTGTAAATTGCAGTTTGTCATAAAGATC TCTATCAGACATTATGCACAAAATGCCAATTTTTGCCCTTGTGATAGCCA CATTGAAGCGGTTGACATTACAAGAGTGTGCTGTTTCAGTAGTTTGTGTG AATATGACATAGTCATATTCAGAACCCTGTGATGAATCAACAGTCTGCGT AGGCAATCCTAAGATTTTTGAAGCTACAGCGTTCTGTGAATTATAAGGTG AGATAAAAACAGCTTTTCTCCAAGCAGGATTGCGTGTAAGAAATTCTCTT ACAACGCCTATTTGAGGTCTGTTGATTGCAGATGAAACATCATGTGTAAT AACACCTTTGGAGAACATTTTGAAGCATTGAGCTGACTTATCCTTGGGTG CTTTTAACTTATTGGCATAAACTAAAGCACTCACAGTGGCCACAATTTCA GCAGGACAACCGCGACAAGTTCCCAGGAACATGTCTGGACCTATTGGTTT CATAAATCTGCCCACTGAATTTAAAAATTCTGGGTCTAGTGGGCCTTTAA ACAACAATGTGCCGGGGGGGGGTGGGTAATTGAACCGGGATCCCCCATATAT TACCTAATGGTTTTGCCCGAAGTTTAACATTTGACACCACCCAAGTCCTA ATTTAGTAGCCCTAAAGAATTTCCTTCAAAAACTACAATGTCACCAGTTG TTTTTGGCGATGGATTTACAGGGCGGAAAAAAAACCGGGCCTAAGGTGAA TTCCCC

>SARS120.CB_C05_-_034.ab1 1600 CATCCTTTTTCTGCTTATTTCGCCCCTTCTTCATACCCTCACACCCGAC ACACCGCGTACACACCGCCTTAGCATTGATAAATAGGAGAAATGAGAG AAGTGCACTACTAACTATAAATTAANCGACCACGACAACGACACACGCAT CGCTCCTTACACCCCACCACCGCGCCCGACGCACTGTCGCGACCCGTCCT CTCTCTCCACGCCCGNTAGCGCAAGATGAAGTACGACGTCGTGGTCAGAC TCCACAAAGATAATCTAGATAGCGCAACAAACCAACATCACCGCATACAT AGACAACCTCGCCATCCCCTAGCATCGAGTCAGACCGCTATCCACGCCGT CCCGGCGGAGTCCCTCGCGTCGACCTCCACCACCGCCCCACCACCCTCG GTCAGCCCCACCGACCTACACTCTCGACCGTCACGATACGAGCAGTACC GAGAGACGCGGACGCCTACTACACGCCCGTACGCCGGTGCGACGGCTGT CAGACAGAGGATGTATAATGATGGGACAGACCGNCTCTGCTAGAGAGCGT GCTATGCGTTATCGAGCGATCAGAATCACCGCGATCTACTTCGTACACAC GACCCAGCACCAGTACGCGACCGACGCGCGACCACCCGACACCACAGC CGACTCCGGTCTGATCGCCGCGCGAATCGCGCGCATCGACCACCTCGGCG TGTCGCAGAACGCGCGATGCCNCGAGGAACACGCGCCCCACTATCGCGCT CGACGGCGGCGACGAGATCCGGTATCGCACGGCTTCGAGTGTCGCTGGC CGGCGAGCGAGTGGGACCTGCGGCGTGCACTCAACTCGCACGACTGAGAC CGCTGACGCGCGACGTCCGCCCCGTCCTGCGAGTCGTCGCGCGTCCGGCG CCGGCCTCGGATGCGCCACGGGCGCACGTGCTGTCACGCTTGCACGAG CGACGCACGCGTGCCCGCACCTGCCCCGCGCCCACGACTCGCAACTCCTG

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>SARS120.CB_D07_-_058.ab1 2247 CAGTAGTATGGATAAGNGAGGAAACCCGATNTATTATNATANTGNTCCNA NTATGACGTCTGATGTNTCATCTCTGCACACAGCTACGACAGAACNGTCT CTACTGCTGCCCTCTNACNCGCTGTCTCNTNCTACTCTGCTCTCAACGCT CNCTCATCAACNTACTCATCNCTACAGAATCAGTNANATGATANGTATCA TCANCAGTCACGCATCACCACNATGATACTGGCTACTCTCTCCTCNACNC TACGCNACAGTACGCAGCACANCGCTACACAGCTCAATACAGCAAACCGC CACACATTACACTACGTNATNATGACGATATGATANCATTCATAGTATAT CATATGTCGATGAGCNGTCGCATGTGCGACGTCCGACACGTGCGACCGTC GTCGTACTGGCGNCTACAGGATNTNAGTCACTGCACGGNCGACTCTCTCN CGACNGTAGCGATCTGTACCGTACGTCATGCGTGCGACCGATATCTGTAN GCATAGGGACAGACGCTGTGNTGAGTGACGANAGCGCTGCACTGCGCGGA CTGCNCGCCGCACGCGAGCGCNGATNGCTCAGTGCGTCTGNTGCTCACAC TCGCTCANCTCACTCGACTCGCTCGCGATACTTAANATATATTATTCGTC GCAGCGTTGCTCTATCTGCTNTNCATATATCTAACTCTAGTCAACATC TATCTGATATCACATCTACGGCTGATANCTGTTATGTACAGCCGACNTCG TCCGCTCATGATACTGNTACGACGCGTGCGTGATANGTAGCAGTACATCG TAGCTGTACTCTGCGTCNACTGAATCGAGNTCNACAAGCACTCGACACTA CACACGTCTGCNCGTCATACGCNTCGANCGTCCGTGCGCTANACTCGTAT GTAGCGATCGTGNGCANCCATGANACGACGTGCGTACTTGACNTCACTCT CATGTCTCGAGTCGATCAGCATGTCAGCATGCTACTGNCGCTATATTGAA CTACGCATACTGTATGGCTACGATGATGGTNCAGAGNGATAGTACGTGCA TCGTACGATACGACGANCNATGGATATCATCAACGACACTATTATAGTGA NCACGACGACACAGAACACACGACGATGTACGCTCAGAAGTACTAGCACT TATCATATCCTCGCGACTACCGTGACTANTGATACATGTAGCCAGATGGT GATCGATACGNATGTACCGTCGCTATGTCACGTCTCGCTCAGTCTGTGAA CTCGTANTCGACNTGACTCATGCACTCGCTGTAGACAGANCTACTTNACA TCGTANCATGCTCATCAGTAGTGTTTCTCTCTTCTACGCTCTTCTGACGA TACATACTCACCGTGTAGCATCTCGCGTAGCTGCTNAGCANCTATTGTCT CCTCTCGTCTCATCTACGTATCGATGGTGCGTCCCATACACATCTGACTG TATGACATCTCGCCCATGCAACACTGCGACCNTCACGCGNACATGTGCCG ACGATATTGCAATCGACGACTAGCAANGTAGTGGACTGNTGATGCGTGAT CTCATAGGTATATCGCGATCGATTATCGCGANGATATTGTTCATGCGAGT NGATCGTGCCTACATCACCACACGTCGCTGCGTACGTATNGTCTCGATGC ACTGATCACTACAGTGNTTACATGCAGGATCGTCGCTGACTNCTCGCACT ACATACTCGACACTCCGGAATGCGTGATATCGATGCGTCGCGTACCACCA GGTCTGTGAGCATGCNACGCCAGGCGACGCTGCGTCTCTAACTNATCGAC GTTGCTGATCGCTGTCACATGAGTGAGTATACTGCGACNATACGCATAGA CGAATACTCTCACGTCGTACTGNCTAGTCTGCGACTCAATAGTCGCGCCT ACGCAGTCGATGTANGTGTTATGTCGTCGTAGCTAGANTGTAGTATGCTG

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>SARS120.CB_E01_-_003.ab1 1337 0 GGTGCGGGGGGTAGGNGAGACANAANAACTNATAACATACTACTACAAT TATATATCATAAACACTCCAATATTATATTCACAATAAAGAATGACTAAC NTGCTCTCGCTGCATCATTTTNATTTCAAACATGATGAAAGAAAGGATAG TAGCAACGNAACTNTNTCGTCACTCTACTACTACACTNCCGACAAAGACG ACACACAGCGAAACAGACGAACGCAACTACTCATCACTCAAAACTACCGA TATCAATATTAATACCTATACCAACTCCATCTCTCGTCTCAGCAACACAC CACAATCAGACACCGACATCACACTTCTATATAATACGAATACCATATCA TGCTCGTACCGCTCATACACACACGCATCACATACTATACGTCACCATCA TATAGCAGCTACGCCANTACTATCATTATATAAATAATAATACACACA TTTATTACAACCATATAGCTGTGAAGACGTTCTACCTCTCTCCATCACAT TCCTTATTCACTACAACTCTACCTATTGACATATTTCCTCACCCCCCCA CTCTCTTTCCGTATATCAACAACACTATCCTCACCCCCCACAACAACCCA CCACAACCCAGGACTTCCCCCAACACCACTCACACTCAATACCTCCATTA TCTACACCAATCATCCCAACATGGAAAAGACCCTCTGACACACGAATTAT

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>SARS120.CB_H06_-_048.ab1 1882 0 ABI GGTAGAGGAGGGGGGGGGGGGGTAGGATGACAGAGCAGATATAACAGAGA TGCGTGGGCGGTTCCAGCTGCAGTCGCGTGCGCGGCTGTGGCGAGTCGCC TATACACCCANTACTACTCATACTCAACGCTCTCGGCTATCTCTACG AGGGCTGTATGAGGAGTGCGCGGAGGTTGTGTCTCGCGTGAGACGCAGCG TGTGGAGCGCGTAGNGAGGGTGATGGTGTGTATCGACATNTATATGTTNG AACGCGATCCTCTATAGATGTACTAGTCATGNAGCGACGCGCTGGNCTAC TGCTCCAGCATATGCTAAGTAGTAGATGTGATATACGAGACATATAGATG CAGAGAGATGATGAGAGAGTATGTACGACGNCTACTAGCGACTCATGTAG AGAGAGAACGGCAGACGAACAGAGATAGTCGACGTGAGGAGAACGGCACA TCTAGAACCTGCAATAGGCCAGACACTACGCGATATGNACGCGCGTGACG ACTGCNGGACGGTCACGCGCTGAGACGCGTCGACTCGTTGCGATGTG TCTGTGATCTGTCCATCCATCTCTCTCTCTCGTCACGTNCTCTGCTGTC TAGCTCTGATGTACTATGTAACGTGATGCTGTAGTTCGTAGCGTCTGTCG ACGCGTGGCTTGCGCGCGTGTGCGCATGAGAGTGTGCGC CGGNAGTGGCTATACCGACGCGACGTCTTCGTCGTCGCCAGCGATGTGAA GTTGTATGGAGCTAGCGACGCGAGTGGACTTGTGAGCGCGTGGCACTGAT GTGGATTACGGAGGTGCTGAGATAGTGATGTCGATGGGTATGTGCTGATA GGTGCGCTGCGGCGATAGCGATCTGACTACGAGGCGTGCCGTGGATGCTT GGTGTGCGGCGCAGACGGTAGGGCTCGTCTGCGATCGGCTGGCGGTGACC GGGCGAGTGCGATCGCAGACGCGCACAGAGTGCGGTGCTGCGGCGTGTCA GTGAGAGTGTTGTCCTAAGACGTACTGATAGATGATGTGAGATCGGACTG ACTAGCCGCTGCGTCCAGAGCGTGATTGTGTATATCAGACTGAGTATGTG CTATTAAACGATGATGNTGGATGGTGCCGTCGTAAGCGACGACCGGGAGC TGCAGGCGCTGCGAGGTTAGGCGGAGCCGGCTGACGCATGTCGTGATCGT CGGTCTCATCGGAGTGGTGCGCATCGTCGACGCGACCATCGTAGTACGGC NAGACAGGCAACATGACCGAGGAGTGGCTCGGTGACGGGTAGAGCTGAGA AGATAAATAGTACACATATCGTCGAGGACACGAACAGGATATCTAAAGTG ACGGTGAGTGCGTGCACNACCCGACGATCTGTATCACAAGGATGACTGCG AAGATGATACGAGATATACGTGCAGAGNCAGCATNGCACGAGTGCNACNG NATGTACTACGCCATCTCATCACATCANATTCGCATACGGGGCGCGNCAT

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>SARS120.CB_A06_-_037.ab1 1861 0 CAACTAAAAAATATGAGATATNTAANACTGCGCATACATAGAGACGAACG ACTCGTTACTACACATAACTACTACATATACATATAGGACCGTCAGAATA CCGTCAGCTNACACGCACTCATCTCGACTACTATATCTTTAATAATGAGG CAATGANAGGGCGATGTAGCGTCTCTTCACACTCGAGCGCCCGCGGCGGT GGTGGNCAGCGACAGCGAGAGAGTCTCTACTACCAGAGTGTTAAGTNTTA GATACTAGTAGTGCTGCATGATGTCGCGCATGACTATAGCAGTGGATG CTCGACTACGTATCGTGACGAGCACGATACGTGAGACTGCGTCGCAGACG AAGCGACGGNTACGCACGCAGCAGCCTGGCGATGCAATGACTACGAGCGA TGTGTGAGTGACACGATACGACAATACTACTCCTCGACGCACTCGACAGT AGTCAATCGTGATGCAGTAGACACACAGCAGTGAACTGTAAGTACACGTA TGCACGATCTGCCAACGAGCACTGAGCACGCAACACGCTCGACGCCAACG ACCAAGACATCCGTGAGCAGACGCACACCGTGCACGACGAGCNACATGCG CAGCTATGTGCAGATGTACACTATCATGTGACGATACGACTGAATGTT ACCATGAGAAGTTCATGATGGATATGATGCATGANACTGATCAGTATGGC ATGATATAGTACGCTGCCGTCGATATACTATCGCACACACGCCGTACGCG ACGGGCGNTGATCAGTGTGCTGAATGTATCATGCACACTGCGAGATGGTG AAATGTCATGATCAAGAAACTCAAATCTGATAGANTCGTAATGCGATGAT AGACGCTCGAGACGCTCATCAATAGCTGATGAGATCATCGAGTTCCGATG TGATTAAGTCTGTGAACACTACCACTAAGTATATATCGGTAATGGAATGT

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>SARS120.CB A08 - 053.ab1 2161 ATAGAGACTACGTTAGTAGTATAAGCAGNACTAGANTAGTAAGTGCATAG CANNTGTGNTATACTTACGACGACACGANTNAGTACCTACGACACGCGAC TAGCGAATGATAGCTNTACATGATGATCTGCNATAGTAGCAGCGAGATAT ACATGTGACGACGTACAGCAAGTAGCAGCGCAGNATATNATAATAGCAGA GTACGANGATAGTTAGAAAGACATAGCTACGTAATGCATAGACGACACAT ATTAGCTATCACTCTTTTATCACGAGCTACNTAGCATGNCGCAGTCATGA NTACGTCGTCATAAACANCGGAGTCGNACTCGCTTTACTAGTACTTCTTC GCTCACGTCANAGTACGGAGATTCATGAGTAGTGCAGNACGTGCCTCGNG TGGTAGATAGGCGTGCNGTGTATCTAGAATACGGCGATGAATACAGATCG TGATGCGTATATCGAATCGCGCATCGCTGCGTTTCGCTCTAGCTACGTCT GTGCGAGACTCGCNCTGCTGCGCCGCGTCAGTCGNCGATCGCGCAGAAGA GAATGACGCAGACGAGACANTGGCGAGTGCTGCCGCACNTCGCTCACCAA CACATCAACNNCAACTANGACGCCACTCCACTATAAGCCACTACTCACTC ACTCTGTGTCTAATATATAATGTATACAAAGAGACATACTGTATATAAAT TATTATATAACAGATGTATATGACGGCTATAATGNTATACTCTCATACAC TCTGCTCACCTCGTCTGTCACACCACATGTGAGTGCGTACATCCGTATAC AGTAGTGAGCTAGAGCCGTAGCATGTCGACACACTTGATAGGTGTACGTC GCGCGCTCATGCTCATAAGCTGATACTGAGAACCATCGAGTATGGAGCAG TGTACTCGTCGCCGACTGAGACACACATCCGCCGCGACGNCACATGACCA CACTCACGCTACACGCCACCGACGGACTGAGAATGAGCATAACGATACAT CAGAATGTCGTCAGAGAACATGCAAGAATACACGCATCACCAGATACCAN CAGAAGTACAGACTAAGACCAATATGGACCAACACACTACTCATANTGCA CCTCACGCATCACGTATGGTATCAGTACGACATGCACGCAGAGCTGCTAT NGACTCTAGCTAAGAGATCAGCATCATATATAGCTGTANCACGATCATGC ATCNTGTATTACTCTATAATACTAACGCTATCACGCGCACCGTTACGATG GCACATAGCACACTAAGCACTATCGATGATACGATACAACGAGCACACGA CAATACATCTGCTACAACAGATCATACTGACAGTATCACACTAATAGACG AGCAGTCGACAAATCTATGTCGAAACGATCGCTGTGCAGGAATGCCTCAG TACGCGCCGGTGCACTACGTAGCTATTCGCTCGTCTATNCAATACAGTCA CAGTCTCATCATGACTACAACACGACACANCCTACCACGCACGCTCACAC GAAGCTCTGACAATAACGAATTAGATCGCTATGAACAGGCTACAGACAAC GATGATAACAGATACGATGCCGCACATACGNCGCCGCCGACACCACTCAC GATACTTCGATATCACGTCGAATACACTATATACAACCTCGATAATCATC GCAATGACTCACGACCGCTATACATACGCGGCATATATCAATGAATACCA GGAGTTATACTCTAACCAACTACACGCCACTCGTCTACCCAACCATATAG CATAACACTNCGNTAGCTATGCACGCCGCGATACGCATCCAGTGTAGNAT ATGGACATGTGCTGACAACCCTCAGTATCCATGAGANGCGAACACTACGA AACACGCGCGTACGCGAGTATCGTCGCTGAGATAACTGACAAATCGCACT CGTATAGTNGTCAAGACTAGAGCATCGATCATCAACTGATGCATGAACAG GATACATCGANCTTTATAGTCTGNGNGTNTATTNGTACGAGCGTAGCAGC GAGCTGTCGTCNTGACGACATGTCGATATCGGATCCCGTGATCATGACGC GGANGTAGACGACATNCGTACGAGATGCACGCTGGATNCGCATGAGTACT GCCGNACTAGTGCTGANGNTCACTATCCTGTCATGCACGCTACTTCNACT

ANTANTANTGN >SARS120.CB_A09_-_065.ab1 1810 0 0 ABI GGCGTGTATATCTGTTGTGTGGCGCTGTCGTGCTAGCTCGTGATGCGCGG AGTTATAGCGGAGCTCTAGTACTCGAGCGCTATAGAGCTGCGTCACGAGC GCTCATGTGCTGTCACGACCTCTCGTGCTGATAACNGACACACAANGTCG CACTCACACATACCAGCAGATGTTAGTGATCGCTCCTCANCGTAACGCTG TGCNGAGCTCGCANCGCGACACNCGANGACAANTNGTATGGTCGTANANA GTCACGTAGTAGTGCGAGCGACATACGCGGACACAGCGCGAGACGACGCG CCTGACAGAGCACACAGTCGACGACGAGTGGTCGAGAGAGCAGCACTCTC TGCGACTNATGCCAGCGACGTGCCGGCTCGCGATCGCTCGCGAGCGTGCT CGCGTCTCGCAGCCGCGGACTCTACCTGCTAGCTAGCTCTGGACTGCTAT CAGATATACATACTATGACATACATACACTTGTTNTTCTCATGTCGNCGC GCTCATTGCCATCCCTNGACGCAGACGTCTACGCCTCTGACAGATGACGT AGAGTCNATCGACGCTGATGTCGCGTCGAATACGCAGTGCTAANATGATA GACGTAGTGTATAGACTGAGTGCGGACAGTNAGACGGGTAGACTGATGTG TCGATGCGAGCTGACGACTGAGTCGTCGATAGCGACTAGTCAGAGACAGC GGACAGCGNCGACGATCTATACGAGGACAGACGATCTGCGTGACTGAGGA TCGATCCAAGATCATGTCATCTGTGCGACGCGATACTCCGCGACACAGNT GTACGGTGAGTAGAGTCGACAGNAGGTAGCAGCGAGAGTAGAGCATGGAC TACAGACTCGAGCTAGNAGTNGCCACGACGACACANCGAATGCATGAGCG CGATGACGTATCGCAGATGATACGAGCATGCTACATCGATCACGTACACC GATGCGCGCTAGTGATGACTACGATGCAGATCGTACGAGCTCCTTATCAG CAGTGTCNCGTCAAAGATCGGNCCTATCGAACTNGCACGCGCGCGCCGAG TACGCACAGCTCGCGACGCACGTAGTCTCACGATGCACTCGACGNCCGCG CATATTAGTTGAGTGCACGCATGAATCGAGAACGATCCGTTCGCACAGCT CGTGCAGCANCGATCAACACAATCGCGAGTGCACATGCGCAAGCATCTCC GTACTCCAGAGAGTGCGCAAGTCACGCACGTGACTCCGTGTGTCNCTACA GCGCGCCGATCCGCGCGTCCTCGATGGCGATGTCTTCGTCTCGCTCATGC CTACTCTAGCACTCACTACTGCGCAGCGGAGTACATGGATGATCAGAACG ATCGGAGCACGCTGATAGCGATAGTTGAGCACACGGAGTATGTCGCGCGC GCGCATTCTACGATCATGTGTCTAGTATGCGGTCGTGTGCACGACGCTGC GACGCGCATATATCATCTTAACGTTGTCGATCGCACGATCGTCGAGCGTC TCGCTGCATCATCTCTCTGATGCTACATGTCCTACGAGCATACATCTCCG ACGATCGCTCAATCCCTGCGACAGACCCGGTAACTCAGATGACGAATAAC ACGTGCGACGCGATGTCAAGGTCGAAGTGGGCATACGAGACGGCGGATCT CGATGTGCGTATTGACGCGGCTCTGGTGGTACGAAGTGATGATACAGTGC ATGTAGTATATGTATGTCTCTGGATGATATGACATGCGATGACATCNC AACATGGGAG

>SARS120.CB_A10_-_069.ab1 1550 ABI AGCCTGGCTGGCCGCGCGAGGAAAGATAGTCAATGAACCACAAAGCAT ATTTATTAGAATAGGAGTCAAGAGAGCACTAGCGACGGTTACGACTCATA TATATATAATAGGGGAGGTGTGAGCGAGATAGTGTTTCAGAGATCAAAAT GCTACAGCAGCGAGAGACAACTCTCCCGGCGCCGCTGTATATTTTGCCTT ATCACTTACACTCTACTACTATCAGAACACACACTGCCCTTACGTCCATA CCCTCCTAACCTCTGTATACACCCCGCTGCGCGCGCGCATGTCATCATATC CTACGACGCCTAATACGATGAATAGAGAGTACAGAGGGATGGACGCAGAG AACACACAAAAAGAGGCGAAGAGAATTGTAGTAGATAGAGATACGTGAGA GGTGGTCTACCAGTTCGAGTATTGGATACATAATATGACATAAGACTGAC GTAACGAGCATGGCGTACGTCGAGGGAAGCTCGAGACGTCGTGTGCGC GAGTCGACATGTACACATGCGTATGTGCTCGCTCTCTGCACCCTGAGATA TCGTTTCATGTGTTCTTACAATCATAATACTAGACTGCGCGGTCGAATCA TATGTACGAATAGAAGAGATTAAACGCGCAGATAGTGATGCTATTTCGCA GCGAGGGAGAGATAAGTTGATAAACAACGCAGTACTGAACGCAATGA GTAGTAGATAAAATCGCAGCGAAAGGTACTTATACGCACAAACACTAAAC CGAGGGTTGGCGCGTAGTGTCCGTACGTGCAGCGAGTCAATCTACCTCAC CTCTGCTGACCTGCATCTTGCAACGTCTAGCGATACGCCACAATAGAGGT TATAGAGCGATAAAGCAAGTATACATCGGACTAAATGAACATACAGATAA CCGCGCTGAGACTACTAATTAGACAGGTCAGTTAGCAGGTAGATGTGAAT GTCTCGCAGGCCTCGCCGTCCCGCCATCATACGACGACACTAGGACTTAT ACCGGCTGCACGACATAACCTGTGTAAGCGAGATTGCTCTATCTTCAANC GGAATGACGTAGATCTAGCGTATTATGAGTGAGAATACAGTGTACATAGT GAGTAGATCGAGAGCAGTACGCAAGATATAGCCATACATTGTGATGACGA

TATAGAAAGATGAGTCCAGATNATAAGCAAGACACGGTACGCTATATACG CAAGAGAAGAATGAGGGAGTATCGAGACGGTGAGACTGAGAGTGTACAAC TGCAGAGAGCGTTCAGCATGCCCTGCGCATACTTAGTACAGACGACTGCT GAATAGCAATCTGAGTGTCGATAGTCATCTCACCTATGCGTAGTCTGATC ATCTATATAACGTCTAGTCAGTCCGCACCACTTATATCACATATCTTACA TATGTCTATGTGTAGTTGACATAGCGTTAATGTGTCATTGATCACAGATA TAGTAGTGTATTGACAATGATGAGACTCAGNAGCTCGTATGTTCACCGCA GAGATNTGTTCGATCTGATGTGGATAGTAGACATCCGCAATCGAGCAACN >SARS120.CB_A11_-_082.ab1 2201 0 AGTCAGTCTGCATCGCTGTCACAGCNGNNTCTGTCCTCNGTNTCAGTTGC TCNGTNNNCNGTCACAGTNTCACGTNGACTCGGCTGTCCGCAGGTAGACT GNGNGGCCGCTTCAGCCGCGCNGNGTNACNGCNGNTCGCGCGACTGTAGT TGGACTGNACNACTGCNGNGCGCTCGNNNATCGNTCGCGTCGTCGNAGTG CCGACGNNGNNATCGTCNGNTCATCGCGNTCGNNCTGTTCNGATGTCGNG NACGCCGACGATCGCGNGCGNGNNTGACGTCGCGACGTGTTCGTNGNGNT TGATGCNGTCGNNGCGNTCGNGTCGACGNTCGNNGTCGTCGTTCGTGTTC TCTTGTACTCGCNGCTCGTCGCACGNTGCGNACGCTNCGNTGTCGCACGT CGGTCGCGNTCGTCGGTGCGCACGCTCGTGCTGCGCGGTCGGGTGCTGGT TGCGTCGTCGTCGTNNCANTCGNTGCGTACGANNTCGTCTCGCGT NACGNCCGTGCGTCGCTGATGTGTCGATNGNACGTGCGGCGATCGCGTGT TCGCGTCGTCGNTGTTTCGGCGCGNTCGATCGNTCGATGTGTCGCGTCGA TACGACGTCGTGTTGCTCGTGCGTCGCGCGCGCNTAGTCGNGTGTCGTG CNTCGNCGCCGCGATCGTCTTTGNTGAGTCGTGTCGCTCGCGGCGCGTAG CGTCGCCGCGCTCGCTGTCGTGNTCTCGCGCGCGCGACGNCGTGCTACGC GTTCGCGGCGCGTGNCGCGTGTGCNGTATGCGTCGTGCGTATCCTNAT GTNTCGTNTCGCGTCGCGTCGCGCGTGTCGGTGGCGTGTGTCGTTCTCGT CGTCCGTGCTCGCGCGCACGNCGTCGCCTGCGCGCTATGTCGTCGTC ATNTCGTCGCGGCTCTCGTTGTCTGTGCGCTTGGCGTGACGTGTCGNCGN TCTGCCGCGTTGTCTCGTTGGNTGCTATGTGCGTCGCGNTGTTGTG TTAGCGCGTCGCTGCGCGTTGTTGTCGNACTCGTGGCGCGCGTGGCGTCG TGCGTCGTGCGTCGGNTGTCGACNTGCGACGNCGTCGCGCGATCTGCCCG CTCGTCCGTTGGCNCGACGGTCGCGTCGTCGCCGCGCGCGCGCGTT GTCGCCTCCGCCGCGNTTGTGCGNTCGCATCGCGTTCGTGGCCGNTCGG CCGGCGGCGCTTGNCTGCTGCNGCGCGCGTGGTGGCTGTNTGCCGTNCGC TGCTGTGCTCGCGTGTCGCCGCGCGTGTCAGCGTCGCGNTCGTG CCGTTTNCGTGTTGCGCGCGCGCGCGCGCGCCGCTCGNCTTCTCCGCTGTA CTCCTTGACGTCGCGCGCGNTCGCGTCGGNATGTCTCTCTGNTTCCTAGC CGGCGTCGCCGTCTGTNCGTCGTCGCCGCGCGNTGTTTCGACCGCGCC GCGCGGTTACGTCCGCCGCTCGNCGTACGCGTTCGGCGCGCGCGCGTATGN CGNCGCGCTACGTTNTCGGTTGTAGCGTCGTACGCCGCNCANCGCGCGGT GGTCGTCTGCTCTNTCGTNTGTCGGTCAGTCTCGCCTGNACGGCGTGCGN NTCGCTCGCTCGTCGTCGTCGCGNNCTGNTCTACGTTCTCGNCCGC TGTGCTGCCGNTGCTCTGCGCGTGTTCNCGCGNTCCTACGNCTGCGAGNC TCGCGCGTCGCGTGTCGCGCNACCGTGTCGCGTCANCGCGNNGTCGNCCG CGTCTCNCTCACTNATTGCGTGCACGTGNCGTCGCNGCTGCGTGTGTGGT NCCTCGCAGGTCTGGNTCTCTGCCGTGACGTCCGTGTCNTCGCGTGNTCG TCGTCGATGTGCGCTCTGTNTGCCGCNCGCGCGTCNCGTTGTGTCCGCGN TGTGTACCTCGNGCGCTGCCGCGGTGCGTGTCGACTGTTGTGTGCNCGTG CGCGTGNTCCGATCTNCGNNCNTCGTNCGTGCCGGTCGTCGACGNTGCTG TNTGTCGCACTTCGCGNATCGTNCGCGCGNTATGGCGCTAGCGGATGCGT NCGTTTGGCCGTGCGNTGTCGNTGCGNTCTCTGTTTCCCTCTCTGCGNGN CTCGCGTCCGNCTGNCTTCNCTCGCCGNTGCGCCGTGCGCCCGCGTNTGN

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>SARS120.CB_B05_-_041.ab1 1010 0 0 AB CTTCTTCTTTTTGAGCTCGGNCACCCCCCACCGCCCCCACCTATGCTTAT ABI AATAATTGAGTACTTGTACTCCCTCATATCGGCGCCAAATGGTGTATGAT TCCTGCAAGAGAACCTCGAACAAGAATTTCCAAAATTCCGAATCCATTGA TGCTAACCCTTGCACTTGCTACAAGCATTGATGCTACGGCGGAGGGATTT ACCTTAATGCTCTTGCATATGAATCCGCTTTTACCCTCCGGCTCGACTCC CCTCCCAACCTCTGGTATCCACCTTTCCACCGCGCCCTTGCAGGAAACCA CACACAAAGCTGATCTTACAGCTACCCGATGGCCCGAGATTGAACTTAAA AATACAACACCCACAACAACACCCATCAAGCCCCTTCACCTTTACAGTCA CAAATCTATAAACGCTAGACCATCCTACCATACCACTTCACAAACTACCT GCACACCCGATCTCCCTCCCAAGACATACCCTCCAATGCAACCGGCACCA CTTCACTACTCAATCATTTCGGCTTACACCAACACAACCCAGCCTCAACA CCCTCCCACCAACCAACCTCACCATCTCTCCCCACCTCTACTCCACGAC CTCTAACATCTCCCCTCTCCTCTCTCCCAGCTTCACCTACCCCCC ACACTCATCAAAGAACATTATCACTCATGTATCTAGACACAAAAACCAAT CATGTCTACGTGATTATAGTATCGTGACTCACCACCCACAGCTCCTGACC ATAATCGACATCCGCCTCCCGCCCGCTCCTATCTAGTCCGCTCAACGCCC CTTATACACCTACTCTCACGCCAATTCTCGTCTGTAATCAGTGCTACTAC CCGGTGAGTCACTGGCTACGTGACCTATTCCTCATCGATCACACATACTA GACAACACTCCCGCCCCCATCTTTGCTTCTCGCCCATACAATCCCTGAC **CCACATCTAG**

SARS120.CB.txt GGACGTGAATGTGTTAATAGAGAGAGAACGATAGCTCAGAACGTAGTGAT AATAGCGCTAGGGGACGNCATGTGTGAGTAGACGCTTNGATCTCGACCAA CAGCTGCATAGATCGATCTAGATGACGCTTACGACAAGGTACGAGTTGGA CCTGACGGCGAGATGAACCGGTGAGCACGACTCAGCCGCCGACGAGTATA GCATTGACGGTGGCGTACGAGTCGATATGATGGAGAGACAGTAATATGCT GTGGTCGATATCTGCAAGGATGGCGAGACAGACAGAATCGTGAACATAGC GAGTAGACGAGTAGTATCGCGTACGATCTGATCTACTATGAATGTAAGTG AGGGATACTATAGTACGAGTATATCACACACAAGTAACAACNAGCCTCAT GCGGCGCGCGTGCGAAGATCAGTGGCATGCGTCGTCAGGCTACGACTCG CGATNGTAATGCGAGCTGCGCCCAGCGACGCACTACTATCNTCGTCACTA TCGANCGACGCCATCACACTACATAACGCTGCACTACCTACTATCGGATC TTACATACGTGCATTCACGCNGATGATAATGGTAGCATATATAACTGTAT GAGATNGTCGCAAGCTCACAACCTACACATGATTANTANTACAANTGAGA TACNTATACATACATCACACGTATCATCGCATCGCTATCTGCCGTCTAAA ACGTCGGATTGTAATCAGAGACCGACATGACATATACACACTAGATGATT ATGCGAGATATGTGGCGATCTACAGGCAGAAGTACACACTTACANTACGA TGCACTACTCACATACGCATTAATGAGGATATACTAACAGTATCACAATG ATACTCGCTGAGACTCGCGTGTAGTACTGCTACATATTGACTATACACTT ATCACATAGATGACACATAAGGCGATTCTGCTAGTATAGCGTGCAGACTA CGTCGTGATATTAATCGCGCACGGATCAGACGATAGTAATA >SARS120.CB_B09_-_073.ab1 2164 TGGTGGGCGTTCCGTGTTGNTCTTCGTNTNGTGNGTNGTCATNTGTCCGC GNCGNGNCGTAGTTCGTCGGCTCTCTCATCTGCTGCTCTNCACTGCGGCT TCGTCGCTCTGCTGCAGTCACTCTGCNTGATNAGTTAGTATGTGN TGNTACTAGTATATGGTGAGTACGCGNCTGCGATACTGATGTCGANTGAG TTGACTAGTAGAGTATGANTAGNCAATGATCTNANATGCGTAGAGCGACT GATCGTGCTTATACTCNGGTCACTGAGNGTCAGACNGACGTANNGCGACT GACGACGCACGTCGCAGAGCGACGAGCNTGCGTAGTACGTAGCTCGNAGA NGTANTATNATNANTAGTAATATNTATAGNAGATGTTACGTCAGTATCGA TCATACTGCTGCTTACTGCGTNACNTNTCATATGTATATCACTCATCG TGNCATCGAGNCTGTCGTCGCAGTATGTAGCATCGNCTGNGTCTNCG TGTATCACTTCTNCNCNCGACCGTCGCGCCAAGTCGCGTCTGGACTCGAG NCAGACNGCGNGTGATNGATCGCACGANGTCGCGCGCGGAGATCCGGCGT GGATGANGANGACGAGANTATCGATGATGTGATGCTACATGATGTAC GAGTCNTTCATGTCAATACGTCGATGATGTGCTCGTCGTAATCTGNNTGA GACGATGANTCTACTATGNTGTAGTGTNGTAGCTGGTCANTGATATCAGT NGNATGTGTANTNTGTAGTGACTAGTAAGTGTGTACACAGCTATGTCGTC GTCGTGCGCCATTCGCGTCGTACGGTAGTCTCGATACGCAGTGAGCGCTG CCTAGTGAACANTGATCTCGAGCTGCAACTGCTCTCCTCACTATCTGCTC ACGTGCCTGCGCTCTCGCATGCTGACTGTTCACTCGTTCTCTACAGC TGCTTCTACGTGTCGCGTGCATCTATTATGCTTAATGATGCTCGTCTGCA TACAGTTACTGCTGANGCTTTGATACGCTATAGTCTGATGATTCTATGTA TGTAGTTACCTCTAGNTGTGCAGTTATGATCGCGTGTCCTCANCACGAGC CGATATCANATCTATGGATTAGCGTACGCGTGACGATAGCACTCACATAG TATCGCGCTCAGTCGCGCACACACTGGACTCGCATGTGGGCTGGCGCAAA CCGGATCAACTACTGACTTAGCATCGCCGCATTCTATCCCACGTGCACTG AGTCCGATGTCGATCAGTCTGCTGAGCGGGCACACGCACAACGGATGGTC GACTCACTCTCGCATATTCACCTACGACGTCTGACGACGTCAGTACACCA GCTCACGTCCGGTGTCAGCCTCAGTGCTGTCGACAACGGTACGCACACTC ATCGACGACGAGCTGCCGCGATCGCGNTGGAACCTACGCGTACTGTATGA

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>SARS120.CB B10 - 077.ab1 1621 CCGAACTGAATCTGTCGCGCTGTCTGAGTAGCAAGACCAAAAAAAGAGAGA CAGTAAACGATCCATCACCCGGAGACGGAGCGGAGTCCGAGCAGCGGACA ACCGTACGACGTTTTCCGAGAGGGGGGGGGGGGGCATGCGACATAAGTCTGTT TTGACGAAATTACCTCATGTCGGTCTTACCAGCATAGATCATTGCTTATA TCTGTCACTGCGACCGGCTACTATATATCAGGCTTTGTGCGGGTGGGCGT GTTTTCTGTGTGAGTTGGATGTGGACTGAAGGGAGAAGTGATNGACGGGG GTGGTTGTGAGCGTGAGGCTGTAGGGCCGGGGAGGCGCCGCGCTGTGCCG CGGAACGAGCGAGAGCAGGTCGGGCGGCGGCAGGTGGTTGGGGCGGATAG ACTGGGATTGTATTGACGTTTTTAACGGTGGCTATGTGTAGACGGGGGNTG AGTGCGCGGAGGCTGTTGTTGGTGAGGTGAGATACGTGTGTAGGAGTTTG TAGAGGCGGCTCGATAGATCGAATCTACGCTACGTGGACTACGTGGTAGC GTCTAGCGTCATGTAATGTGGTGAGTAATTACGTATTATACCAGATCGCG TACTACAACATCATTTTGATTCGCGTGGACACGTAGAGTGGCGATGATGG GATGGTGTAGTGATGCGGTCAGAGCTCAACAGTTGCCAATTGTGCGTGTC TGGCGAGCGGAGAGACAGTTAGGTTGAGGTTCGCGCGAGATGGTACGTG TGCGTGGNGTNGGTGGTGCGGCGCAGCGACCTGGTGAGTGTGTGCTGGNC CGTCTGGATATCGCGGCAGGGAGACGGACTGTGACGCGTGATGAGGGAAG AGTCGCGTGGAGNCGTGTGGCGGTGTTACGNGTGTGNACATCACGGNTGG TGTCGGTGAACTGAAGGTACTCGCGTGGTGTTGTGACGTGACCCGAGGTG CAGAGCAGCGCTACTGTCAGGCGCTGCGTCTACCAGTAGGGTTGTGTAGT GCGGCGCCAACTGCATCAGTGATGTGGCATGGCGTGGTGTGCGTGGAAGC GCGTGCACCCGGGTAGAGGAGTGCGTAGCNTGGAGACCNGTGCGAGGTGG AGGAGTGCGTTTTGGATGTGTCGGGGTGCGCTGCGCGCGTTGGAGCGAG GACAGNTTGTGATCATCTGACTCTGACTTAGACTGCGCAGATGTNGTGCT ATATGGAGTTTGCAGCGGTGGGTGTGAGGGTAGGTTATCTCTATATATCC GCTCACCGGAGATTATGGCTTCATTATCGGCGCGTGGTCGGGCAGGATCG GTTGAGTATCNAGGTGTGTGTTCGTGCGTGCTGTAGCTCCTCGTGCTGTG TGCGCGCTGATGCGCACGGGCGAGATGTGCGCGCGCGCGNACNTGCCNGC CTATNACTCGCGACTGCGCGTGTTCTGTGTNAGTATAGCTGGGGTGCGCC GTNGNTGTCTACCGTGACTACGCCGTGAGATGCGNTGNGATGCGCGGTCC ACGCGCGTNTTCATATGNGTGTTTGTGCTGTGTCGTGTGACTTCTGTTTC GTTGCGTGAGATGGTCGTGGGCCGGCGCGCGCGCTGTGATGCTANATCCTG AGGCGCGCTGATATGGGGATA

>SARS120.CB_B11_-_090.ab1 1468 TCTACTCTTGCGTACTTGCTCCTAGTCCTCGCTGTTGGCGACGGGCTCGT CGNCGCCCTCGCCGCCGCTCTGCTGCCTTCCTGCCGGACGATCCGATTG TCATAAGTCGCACGGCGCGCGCGCGCTGCGTTGCTTCTCGNCACGACGC GTCGTTGGTGCGCNTCGGCTTCTGTGCTTCTTTACACCTGCGTAGCTCAG GNTGTCTGCGGCTATGCGGCGCGCGCCAGNTCCGTGCGGCGCGCGCGCGC CCGCCGGGCGCTTATCGCCCCTCTGTGCNCGTCGCCGCCTACGTGCGCTC ACNTGTGCGATGTGCGCGTGCGCTGTGTGTGCTTCTGATCAGTGGCCTC TGTGCTGTTGCTGGTGGCGCCTCGTCGGGCCGTCGTGGCTCTGTTCCTCG CNCGTGTGCGCTGTCGTCGTANCCTGCGTGCGCCTGATGCGCGTTC TTCGCCTGCACGCTCTCCGTTCGCTTCTCTCCTACTCTCCTATCTCNTCG GCGCGACGTGGGCGCTGTGGTGCGTAGTTCGCTGTACGTCTCTGTGCTGC GGCTGGTCTGCCTGTCTGGCTTTTCTGTGTCCTCGGTGCCCTTGTC GNTGCTACGACGCGCTTCCGCGTAGGCCGCCGCGCGCATCGTCGCGTC GCGCGCTCTGCGACCGTCTCTGCCTCCGTTGTGGCTTCGTCTCGCCTGTC TTACCCGCTCGGCTCCGCCTCTTGCGTGGCTCGCGGTGATCTGCCTACCC TGTCGTCTGNCNTCGCGTCGTGCTCGCACGCGCGCGCGCCCTCCGTGTGAC GTGGGCCGGAGACCTCNGCGGCTNGCTGTGATGTCCGTGCTCGCTTCTCG CGCTCTCTGTCGTCGCCGCCTCATGCTGCCGGTCGTCGTCTTC GCTCTCTGCNCTCTCGCTCCGCCCGTCTCCCACTCCTTCCTTCTGACGCA CGCTTGTCTCTCTGCNCTCTGCGCGCGCTCGCTCGCGGTGCCGGCTCC TCTGCGTNTCGCGTCCTCCTTCCTCCTGCGCATCGTCTGCTGCCACNGCG CGGCGACTCGCTTCTGCGGCCCTCGCACCGGCTTCTTGCACGCNTACGTG CGTGGCACCCTCCGGCGCTGGGCTCCCTTGGTTCCTTGTCCGGGACCTCT CGCTGCTTCTGCTTGATGATCGTGGTTCCCGTCGCTGTGCTCTCTGCTGT

>SARS120.CB_B12_-_101.ab1 1828 GATTTACATTAAAATATTCTATTCTTAACTATGATCTCAGTATCCGCCGA CGCTGCACATATATCAGCACCACGACCACAACCCCCAGAACACGCGATAC ACATATACAGCAACACGCAGACACATTCATAGNTCATCACGTCTAGCCGT GTTTGGATGCCGTCCTGATGCTCGCGCGCTCGACTACTTGCATGAGG ATACACGCATGGCTGCCAGCGGTGANCGCAACCCTACAGCTGCGTCCACG CNACTAGATGATACCAGCGACANCTTGAGCAGCGATGTGACGCTGTCAGT GATGGCCTCTGCTGACAACAATCACGAGACCGTAGTAATCCCTGCTCACG ATCAACTTATCGCTCATTGTATCTTCCATCTTNTCGTCGATCTCGTGTCT CCTATGCTGTACTCTCATATGCGATCAGATAGACAGCGACTATACTATAA TACCACGTACGAGACTAGGAACAGACACACAGATCGCCAACACACGGAAG TGTTCATGAACTATAACGGAATCTGNAGCGTAGACGTCTGAACAGTTAAG CAACACGAAGACGGACGTGATCATACAGCGAATGCAGGTCAAGTGTAGTG CGAACGACGACGCGTATACGACGCGTGTCATCACACACTAACGATCGCG CTTCTGAGCTGTCGATACGATAGTACGACTGAATATTGCCTAGTAGCGCG CGTGCGTATCGCTTGCATATGTGTCCTTGAGCGACGTCTCTACGCTCGGC TCTCTGCTATACTACGTGCCCAGCNCATCGAGGCTCGTAGCGCGTCCGCT ATGACTCTGCAGGGCTACGTGTATTGTCGGAGTTGGCGGCCTCNGCACCG CGTTGTGCGAGGCGATGAGTATACGGATACCTAAGGTAGATGTCAGCACG TATGACATATGGACGACGTGCAGCACGTCTTGGTAGTACCACAGCACGAC GACACATCTGCCGAGCGATACTATAAGTGTAGAACTCTCAGCAAGAATGG CGGCTCATGCGGAGACCGTACTAGTACGGTATGATGAGGTCGCCTGTAGA TGNACGACGCGATGCACACTGACACTCGCTAGCGCTATGTACTGTGACGT GACTACCGCGACGTCGTACACTCTTCTGTGACATCAGCGTGGAGACTAAA ACGCGCAGTCGAGATAGACTTGTAGATAAACATACGTAGAATGCATGATC ACTCTCGACGTGCAGCCANTCGTAGATATCCGAGGTGCTGGACAGACGTG GACGTCCACGTGAGTACACTCGCATATGCGTCATGTNGGTCACTCNCTGC GTTGTGTCTCCTACCTGANACGCTGTCTGGCNTAGCGTCATGATATAGTC GAGTCTCGATCCACTCGCGAGCTGCACAATACGATAGTCATGTGTTGTGT CACTGACGCGCTCATATAGTACACGAGTATCTGACGTGTNTGCTGTCANA ATAGCCACTGCTGTCGACGTGACTCTGNAGNAGCGTGTGTCGACTGTANT GTGGTACATACTGCTGTCTACCGATGTNGCACAGATCTGCGGTAACTCGC GCNTGTCAACTAATGCTGAGTNANCGACGCGCAGTCGACGTCATNCACGT CGCATGTGTGACTGTCGTGCTGATAACGCTCATCGTCAGCCAANTCCTNT GACTCGACACTCGCTTGACACANGAGACTCGCTCTACACTGNAGATCGTC TATACTACATGCTGTTGTTCGCANTCTCAGATGCTAGCATCANGCTGTCA **TGTCACTGTGGTATGCATGCGTCGGATN**

>SARS120.CB_C01_-_002.ab1 1435 AGGGGAGGTGNTAAAGGTGAGCACGAGNAACGAGAGAGCGCGAGGACGCA GTGTAATAACACCTACACNTCAAAGCAACGGGCGCCTCCCTGTATGGCGT AGGGCATATGCGTTGCAGCGGCGGACACGAAGCCAATGGTACCAAAAAAA GGGGGGGAGNGAGGAATGGGGGCTGCCTTTTTCCCTCTTAGACTACGCAA AACATACCCACATTGGCTGCTAACTGTACTGCAGCACCGACTTGTCTCAC ACTGTTTACGCCTAACGGAGGTGATGCGGCTAGGAATGTAAGTGGGAGAT GCGTACATGCTAGTGGGCGGCATCCACGTATACTGTCTACAACCCACGGT TGAGTAACACTCGATCACGGAGTGTATAGCGTAGCAGACCTGAATCTATG TCCGAAAGTAGGCGGTCCTGTATACATCGTCTATGATCGACAGCTGTGCT TGACACGGCGCAGATAGTGCAAGATGTCATGCTTTCTATGTCCGAACGTG AGAGGTGGGAGAATGAAAGGATGAGCGCGGGGGATCCGGAGGATACTTCCG CCACGCCTCAATGTCGCGTGCCGAGCACGAGCACGNCGACGATGTTGCTG ATAGTGAGACACTGCAGTGCCAGTCTGGAGCGCGCGTGAATATCAACGCG CGCGCGATCGCTGAGTGCTAGCGCACGGTNGGAGAGCTGCAGTGAAGAGT AGAAGTCGACAGGTACTCGATACCGCGCACGTCTAGCGACAGTGCAGCNG TGATCGAAGAGTCTGCTCACTGCGCAGTCTGACACACTGACGCGCGT AAGATAACTATCGCATGACTCACNANCACGCTCTCAGCAGCGTGACTGGT CCACTGATCTACCACATGCGAGACTACACGAGACGCTGGCGCGACTGCTA AGGTGCGTACATTACGCGACGCACCGCGGTGCCGGCGCGAGCGGTGAGGA GTGGGACGGCCGCTGAATTGAGCGAGCCTATGCTCGAACAAGACTCC

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>SARS120.CB_C07_-_050.ab1 2170 AGTTATAATTTGGTGGTAGTTGTAGGTAGTNGTGTGTGCGACTGTGTNAC TTNTNNANAACCACATCACCACACACAGCGTATGTACTATCTTATATCT ATTCCTACTNNCGATNTCTATCTCTATCGTACATAGTCATCTCAGTNACC TANTATCACAACATTACACATCACTATCTTATCTTANTCTACTGTANTACT ACAGATAGACAAATAANCCTNCACCTGAGATNAATCTCAACATGAGACAT ATATTCTGTTACACACACATCTACGTCCCTCTATCTCTCATCTNCTTNTN ACTAGCATATATGNANCANCACGTGTCAGGTGACAGAGGAACANACGCGA TCTNAGATGTNANTGACGGACAGTAGAGTCNATATGCTGTATCAACTGTA NTGTGACGTGCAGCATACACACACACCAGAGGAANAGTCTCTAACGCAG NTGCGTAGTATCTAGCATCGTTACTGACACACGANANGACGATGACATAC GATANCTGAGATGATCATATCAATGCACGTATGATATAGCTGATGCGAGC GACGCGTGAATCACAACGAGATATCGACACGCAGACTTAACGCGCATGTG CTGNGACGTCTACGACGGTGACTGTAGGCACGCACTCTGCGCCNCTCACT ACATGACAGGTTATTATCCTCTATCTTTATAGTTATATATCTNTACTTAT NTTGTTCATACTATGCTCTACGAGACTATATACTGTGTGCGAGCGGATAG CTGACTACTGCTGTGTGTGCGGCATGTCATCAATCATAACTCTATGTG TGTCGTATCGTGTCTCGCGTGCTTGTCGCTTATACATTAATACTCTACTG NTACGATANATGTTCTCTCGACGTGACTCTGACAGAGCTATGTATTACTG CTTCTCATGTTGTCTCCGTGAGCATATGCGTTGCACGTTACGTCTTGACT GCATCTTTATATCGCTGTGCGTGTCTCTGACTCANATTGTTCTTTGCGTG GTTGTCAGTACTGCAGTCACGTACAGCGTATCTCTGTGGCTATGGTATNG NTATATAGCTGGAGCNTNCAAGTCGATACNGATGTCATGCGTCGTGTCGC GTGAGTGCTCACTCTGCTTCNGCAGACGTCGTCGCTATGCTCACTCTATA TACTGCTGTTGCTGACGCTAGACTCTGTGACAGTACTGCTCTGCTCAGCG TGTGATTGAGTGCAGTGCNTATGTAGAGNCGCGTGANTGTACGACTGGTC CGCTATAGCGTCTGTTACTCGATTGCATCTACTCNACGCGTGCGTCGTCN TACGACAGTGTGTACGTAGTGATATCGTATGATCGGGAGTATGATGCNTC TGACTGCGCGATGTACATAGGGTCGTTGCTGTGTNCTCTCTCGCGCGTCG TCGCCTAGATCGCGCACAGCCTACTCGCGACTCATACTCGATCTTCTGGT ACCTGCGCGTAGATCTTCTGCTGCAATGATAGTATTCTTCTATATGTCTT GGTCTCGAGCTGTCGGTCAGAGACTTGATNGCCGTCCTGATGTAACGTTC GTACGCATTATCGTNCGCATTGATGCACAATCTNAAGCTATACCGTACAC GANTGTGCGTCTGACTCGANTNTACGCGGCCGGACTATAGCAGCAGCGTC

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>SARS120.CB_C09_-_066.ab1 1565 GTGGTGTTTGTTGATTGTCGGTTGTGTCGGCTTCTATGCGTCTTCGTC GTGCGTTGAGTGCTCTATCTANTGTGATTCTCTTCGTATATGGTCCCTTC TGCTGCTCTGCTGTGCTATGTGCGCTTGCATTGTACCGCGTCATTAGTGT TNGTTGTTGTGTAGTGACTCGTACTCGCGGTTATCTCGGTCTGTACTGTG TCTTTCTGTACGCTGTATATTTCTATTATCTTCACTTTTATTGATCANAT ACTACAAGCGAGCACCTACAAATCGCATACGACTACTACCTGCCCGCTCC CGCCTTCCANACTTCATGAGCAGCATCGTGCACTGATCTCGACGCGCGCT TCTGCTGCGTGTTCTTTGTGATACGTATGTGCTCTGGGCTTGTCTCTCGC ATACTCTGTCTCGTCTGAGCGTCCCACGCTCTCGCCACTAGACAGGTCGA ATGCGTCACTGTTGTCACCTGGCTTATACTTTAGACTATTACGCCACG CCATCGCTGCAGCGATCTAACTGTTACGAGCAGTGTGCGATAGAATTGAT CGTGAGATCAAGACATAACAATTCGATCGTGGTTGTGGGTATCAATTTAC TGTCTACTGATATCGAGTGCTGATGCTCGCGCTGTGTCTTTGCAGACTGC TCATGATCANCGCGAGCTCTTGTGCCTCGTTGCTTCTAGGATCTGTATTC ACTCTCGTGCCTGTGACGTGCGTCTGTCTCCGAATTGAGCACTATTGACG TATTACCATCGCCGCTTCGCTTGTCTACGTCAGTGCTGTCTCGTGTTTTA TGATCCATCTCATCGCGCATGTGTGACTTCTCGTCGTCAGCTCGTCGT CTTTCCGTTGCGTACGTATCGTCTGCACGTGAGATGATCTAGTGAGGTTC GAGTCAGAGACACGTCGACTTGCGNCATATCGTATACGTATGAGAACACT GCGAGTGCGCTGATGTCTTGCTAGATGCTCCTAAGTGTTGCGGTTCT ATCTAGTATAGTTCGCTATACGAGCCACTCGACGCTAGCGTGGTAGCTAC TCACAATGATGATTATAGGTTGCATGCCGTCCAAATGTGATATACGCTCG TCTNGTGCCGACGTCTATGACTATGTTGGGTGCTGTCGACATATATCATA TATTGCTGTGATAGCTCGTATCCTGAGCGTGGATCTACCGTTCTAGTGCC GCTGATACTTCCACTTCCGACTCTCTATGTCTCACGTGCTGCTTAATGTC TTCTCGTNCTCTTGTATTTTCGTCTAATTCTCGTCTGTGTGTCTCTGCTT TGGTGTGTGATACTGCCTCTGTATACTATCAGTTCGATCAATAGCTACAA CAGACTATGATACTATATATATCTACACGCTCCCGACTAGTCTATGTAT ACTAGAAGATTCGATTATAGTACATCTTCTCTAGTGTTATTGTAAGCCAT GCTTGTACGTCGCATNTTCTATCGTCTCTTGCTNTGTCTGATCATGAGCG **TACTTGTCAATATGN**

>SARS120.CB_C10_-_070.ab1 1338 GAGTGTGAGTTGGAGGTGTGCGTGGCGATAATGTAGGTCAGGATAGGT AGTCTACGCGATGATTTTATGGGTCTTCGATCCCTACATAGAGTCATTGA TAGCACGACAAGCACGACGTCATACATGAGAGTATATTACATCATGTGGT GTGCCGCAGACGGAGGGCTCTTGTGTTCCGTGTGTATCCACGCCTGAACG CGCAGAACCCTCCCCGACCTCGCCCACCCCGCCTGCTCCCATCGCCTCA GCAGCCTTTCGATCGCCTACGCACAGCCCATCAGGTGATAGAAGGAGATG TAGAACGTAGAGGCGATAAGAATAGTGGAATAAGTNCCCCTCGTAGAGTG GCGATGACGACTCTCATATGATCACGCAGAGCCTGAGGATTTCGTCCGTA CTCAATCTACTCTAGATCTGAGCATTGCGAATCGAGTTATATTCTGCATG TTGGACCTCGCTCTCATAATAGTCTACGTGTTATATATACAGTGGTGGAT GGAGTATCTATAGAGGAGATGTAGGAGGAGTGGTGACATACAAGAAGAGC GCGGCGACTACTAGTACGGTGATCGATTGAACTCTCTAAGCTAGCAGCGT AGGCTGTCGGAGCGCACACCCAACGTATACACCCGAGCGGTTATGTTTGT TATACTATATCGTTCAACACTGGGTGTGATCGGATTCCGAAAAGTCATGA CTCGTATAACTTTCCTTGTCAGCACTCGTGGACGCAGCGACGGTTACCAT GTATACTATATGCTGTTATGAATGATCGTCGAGTACCTACTCACCTATAG AGCAGAATAGCTACCACGACTTCAAGCATGATATTAGCGACGTACCTCAC CCACGACTGCTGATTGAGTAATACTAGTGCATCGGCTCACGTCGCCATTA CGCGACGCGCGTGCACACTGTTGATGACTCTGATGCACTAAGATATTACT

GATCCACTGCGCGACTCGTGATGACGTGATACACTACATATATCTAGAGT TCGATTGTANTGCGTTGTATGATTGTGTGGTAGTGGAAGGTGGAGTAGAC GATGTACAGATATATCCTGCAATATTAGTTTCTCTGGTGATCGGTCGTGT ATCGGACTGTACACTCACTGCGTAACCAGTGTGGAGAAGAACAATGTATC TGCTCAGCCTCCCGTATGATGCTGAGGATAATAGATGGTGCTGCCTCTAG TCGATGTCTAGTCCCTAGAATNATACTCTGTGATGGTGGCGTGTGTATCG TCTGACTTGTGTATACGTGGTCATTGGTCGCGCGAGGAGNTGAGCGTGNC GATGTAGATGTGCGCAGAACCGATACGATGCTGCGTCG >SARS120.CB_C11_-_084.ab1 1749 CGATCGACGCGACGACTCGTCGCTCGTNACACNANCACGTCAGTGGTCGA CGCTCGTGTGATGNTGTCGCGACACGNTACGTCGCGAGCACGCGCTGCGA GACGACGCGCACACTGCACGCGCACGNACGAGACGTAGACACGCAGTNAC GNATCACTTAGGACGACNCGAGANACGACGCACAGCTGTTTTTTGNCGA CGACAAGTCACGACGCACGGCACCNACCTCGTCTGTCAGTGCTGTGTGTC GTCTCACCTGCGCGTCGGCGCGCGCGCGCGCCCATGCGTNCGCTACGG TGTCAGCTGAGCTGTAGCTCGCTGTGGCGTGCGCGCTAGCGTGCATG AGTCTGTTATTATGACCTTGATATGTACTNTTTTTGCGNTGTGAGCTGCT CTCGCTCGATGCCATGTGATGCGCATTGNGATCTGTCGCTGTTGTCGTGC TGTGTGGTGACGTCGAGATGCTGTTGGTCGNCGGGCGCAGTGCGCTCTGA CCGTGAGTCGCGGTGGTGTGATGTGACTGACGAGGTGAATCTCATGAC TTTCTGAATGATTTGTGCGAGTNATGTTTGTAGTCATCTGCTCTGTTGTG CGGGTGACGCGTGTATTGAGTGTGTTTTTTGAGCTAGTCGTATGCGT CGATCGTGACCGNCGTGGTGGCTGCATGCGTGCCGAGAGGGTAGTCGGCT CGTGTGCGCGCGCGTGCGTCGGTGAGTCNAGCTGAAATCGGAGATGTCGT AGAGTAATATGTTTGAGTCTGCCTGTGTCTTGATTTTGTGTTTCGTGCTG AGTGTTCAGTNTCGCCCTCNNCCTGCTGCTCGTGGATCGTGCGGCTTCGT AGTGTCGCGCGCCGCGATCACGTGTGAAGGTGAGTTCACGAAGNCGTCTG AGCCTGATGTGGATGCGACCTGCNACNAGTAGCGCGATGAGCGTCACCGC GCCCACTGCTCGACGCAGCCGCCCTCAGTTACNCGTTGCNCGTCACGAGG CGTACCTGTNCCGCTGCTCGCCTGCTGCCGCGNCCGACNTCTGGATGGNC GCGGTCCGACGATGTGACGACAGCGCTCGCCTGANCGGACGGCGCGTGTG CGCGCCGCTGGCTGNACATCGTCGTCGCCCTGTCTGGCTCGCACTGCGAT GTAGTAGATGTGGCAGCNCGACTGTAGCTCCACTCGNNCAGCGTCGTGCC GNNACTCGCGTCGTAGAGTCTGGACGATGNTGTTGTATGTGCGTGATCGT AGTGCTGCATGTGGTGCTCGGTCATCCGTNCTCGTGAGCGCAGCGATGAG NAGCGCATGCTGGTGGCTTACAGNTGTGTGCTGTGNAGTATATCTCTGAT GCGTGTCGCCGCTNTGTTCATCTATCTGTGGCGTTGTGGNNCTAGAGCGT CATGACTGTGCCGNCGGTGCGTGACGTCGNTGTGCGTCTTCNCGCGTCTA TACNCGNCACCGTCCGCGNCGTGTCGNACGTGCGTGTGATGTNACTAGTA CTGCGCTGCGTGNCGCTGCGGTCGNTGTANAGAGCGNCGGANNATGATCG CGCGNTTNNGATNCTCNACTGACGCCGNCTGTNTCCNTGCGTGGTTGNAC GTGCGTGTTCCTGNCGNNCTGGCGCGCNACCCGTCGATGTGGCTGCGTCT GTGCGTTGTCNTGNNTGCGCGNATGCGCTCTACGCGAGNCGNCGTGTGCA GACGGNCNTGACATCGCGNCGTCNCGACGCTNCANAGATNACNNCGACN >SARS120.CB_C12_-_098.ab1 1279 0 GAGGATGAAGGTTAAGAGTGATATAAGAGGGAGAGGGCGGGTGGGCGCGG GTTGCATTTGTTCTGTTTGTGTCTGTTTGTTATAATTATCGAAATAA AATAATTTTTTACGCTGCGGAACGCGACGAGGCGAGACGTGTTCTCCT TTTGCGCTTGAGCGCCCCCAGCGCACGTATCGCGGCACACGAGCGCGAG GCGCCTCACATCCCCGCCACGTATGACGAGTAGAAGACAAAAGTGAAGAT GAGTTCGCATAGGACTATAAAGGAGGCTTCCAGCGCAGCATCGTGCCGTC AGCTGCGGGTTGTTGTGTTCTCGTGACGCGTCAGGTATGATATCGGCGAA CGCGGGGAGGTGTGGACGCAGTGGTGAGAGCTAGTGGAACGAGTATGGA GTGTGCAGGAGGATGGTAGCTTTCGCCTCCGGACGTCGTAGCGGACGA AAATTGGCTGTGCAAGAGCTAAAGTTGTCACTGCGCATTCTGTGAAGATG GTAACACAGTATGATTAAGGTGTAATATGCTAACTTCATGATATACAGTG TCTAGCGCATACCTGATACACACGACGAGATTGCTCATGCTGGTGCGTGA CCTGACAAGAGCCTATCATTATGTTGGTTATCTGTTTGACTTCTGACGTC AGCGGTAGGCTAGTGAGGGTGTTAGTGCCGAGGCGCCAGCGCGCGTGGCG GCCGCGGAACGGATGGCGCGTGTATGTTGTAAGATCCACTCACAATAAA CGCGGGTGGTTGATGTGAATATAAATTTGATGTAGCGTCCGATACGACGA AGGGTAGGCGCTAAGGGTTTCGCAGTATCCACCCGTGTCGGTGAACAACT TAGACTATATGATAGCGACATGCAGCTATATCAGCATCGTTCGGCGACCT

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>SARS120.CB_D02_-_014.ab1 1015 ATGGGGGGAATAAAAAAGAACAACAGAAAGAAAAAAAAACCACCTTCCCG CCCGCGCGGGAGAAGGGAGGGGGGGGAGAATTTCCAAAAAACAGGCA GAGGGGGATTTTTCCCAGAAAGAAAGGGAAAAGGGAGAGTACGCCACGCG GAACACGGGACCCCTTTGCCCCCAACAGGGAGCGAGCAGCGCACGGGGCG AGGACACAGAACGGAACGCAAAAGACAGCAAAAAGCCAAGCAGGCG AAAGACAACCACACCACCAGACGGACGCGGAGAGAGAAAGGGGGAGACC GAAACCGCCAGCAGAACACGGCACAACAGGAGGCGACGGAGAAGGGAAAG ACGCCAAACACAGGTCGGCACAACAGATACGAAGAGGCACGCGACAGAG GAGCAGACCCGGACGACGGCGGGGGGGGCCACGCGCAAGAACAG AGAGAGGAAGCACACGGCAGCGCAGACAGTGCGAGCGCGGACGAAGGAGA ACAGGACGCACAGAACGGCACACACGCGCGGGACGGGACACAGGGACACG AGGGAGCAGGCAGACGACACAAGCGGCGAGAGCGCGGGCGCGAGCG CAGGTACCCAGCGCGACGCACGGGGAACGCAACCTCGCCGCGCGAAAAGG GAGAGAACCGCGCATGGCGCGTGGCGCCCCGAGCTGCAGCGCGAGCTCG AGAGGTGAAAGCTGGAGCGGGGCGAGACCCGGACGCGCACCGCCGCTA CGACGCACAACCGGCGCACGGGAGAAGTGCAGGTAAGGAAAGCAGGGCAA GAGGACGTCGAACGC

>SARS120.CB_D03_-_026.ab1 829 37 453 AB: ACTCGTCATATCGGTGCCTGTGTAGGATGAAGTTCCTAATGATCTTACTA ABI CAGTCCTTAAAAAGTCCAGTTACATTTTCTGCTTGTAATGTAGCCACATT GCGACGTGGTATTTCTAGACTTGTAAATTGCAGTTTGTCATAAAGATCTC TATCAGACATTATGCACAAAATGCCAATTTTTGCCCTTGTGATAGCCACA TTGAAGCGGTTGACATTACAAGAGTGTGCTGTTTCAGTAGTTTGTGTGAA TATGACATAGTCATATTCAGAACCCTGTGATGAATCAACAGTCTGCGTAG GCAATCCTAAGATTTTTGAAGCTACAGCGTTCTGTGAATTATAAGGTGAG ATAAAAACAGCTTTTCTCCAAGCAGGATTGCGTGTAAGAAATTCTCTTAC AACGCCTATTTGAGGTCTGTTGATTGCAGATGAAACATCATGTGTAATAA CACCTTTGTAGAACATTTTGAAGCATTGAGCTGACTTATCCTTGTGTGCT TTTAGCTTATTGTCATAAACTAAAGCACTCACAGTGTCAACAATTTCAGC AGGACAACGGCGACCAGTTCCAAGGAACATGTCTGGACCTATTGTTTTCA TAAGTCTGCACACTGAATTAAAATATTCTGGGGTCTAGTGTGCCTTTAGT CACCAATGTGCGGGGGGGCTGGTAATTGAGCCAGATCCCCCATATAGAAC GTAATGGTTTGCACCAAAGTCTAGCAATTGCCACACCTCAAGCCTAAATT AGTACCATAGAGAATTCCTCCAAAACTACCATGTCCCACATTGGTTTTGG **GCAATGCTTTTCACGCGAAAACAAACTGT**

TGAGTAAGGATATAGTAGATGCGTGAAGACTGCGATGAGCTTGGAGCGAT GAGATAGACGATGTCGNGCAGCGACGATANTCGAGACAGACTACAGTAGG AGTATACGANCGTACATGCGAGTAANCGACTCACCTGTGATGTCGACA CGTGACNAGACGTCACGTAATGACTGTATCTGCTGACGTCAACAGACGCG AACTCGCTGTCGTCTCCGAGACCGAAGNTGAGCGCGTCACCGAGGATCGT ATGTCATGCTCTGCGACTAGAGTGNGCTGAGAGACTAGTGATCGACGCTG AGCATGATTGCGACGTNCGGAGTATANGACATACGATATAGCTAAGGNTC GATACCATCGCGTAGCGTAGCTATGTAGNACGTGCAGTGANCATCGTTCT ACAGCTNACGTATCGATCAGCNCGTGCACGAGCGTACTGCGATCTGANAG CTGCATGATGCTAGTAGCGAATGACACGAGTATACTGACTCAGATACGAT GTTGAGTCCANTATGATGCGTCAGTATAGTACTAACATCGTATATGTACA TGTGATCTCACGACGATGCTATGTGCTGATGCTAGGAAACNGTGNTGCAC GANGNCTGACGCTCATGACGACTCGACTGTGCGCATGACTGTGTCGCA NGCACNACTACCGACGACGCGTATCNCTGGCACTGANTGCATCAGCCGA TTATCACATGCGACGACAGNCCTCTGCGTACGGCTNCGCATATCTGCAAT ACATCGCATTCACGCATCGTTCGTNGCATGCTGTCTGCACACCGTGCTAC GCTACNGNGACACNCGTGCTGTAGCAGCGCTAGGCGNCATGATNNACGCN TAGCAGCGTACGACAACNACAGCGTAGACTGACGACTCNANCTAGACTCG ATCGTGAGATCTGNGTCACGCNCGACGCATATAGACGATCGGCATGTATG CAGTGATGTATATGAGACGNCNANTGGTCGATCACGGACGACTGNATACG CGAGCTGCAGTTGACCGCGTGATGGACGACACACACATCTGTGATAACGA TGCGATGACGCGNGATCAAGACGTCATATGTAGGCTGTACTACTGANCCN TGTAGACACAGCTGGACGACGTATGCNACTGACATTGTATATTGACGTAT AGCTCATAATANTGACGTGAGCTAANTAACTAGATGAGTGGAAAATCGAT ANGTNGATCGACTANGACAGTCGATCACAGTCATGATNACTGCGCAGCAC GTACTANGTAGTCGCGTCTAACTNAGCATATATGTCTATANTGCTTGATT GCGTATGAGCATTATAGCTATGATGATNCAGATCAGCGCTAGCGACTGAC GAGACTCGATGTNGAGACNATCGATTACGCACTATCAGTAGCTGCAGCTC **GNANTCGTAGACGCGTGATGACATCCGN**

>SARS120.CB_D06_-_046.ab1 1437 0 AGAAATATCTGCTTTNTTNAGTTCGGGCCAGACGCTAGGCCCAACGGGCG TGCGGAAGAAACAATATGATAATTCCGNGCGAGCCGCTCCACAAGCGAC GCTAGGGCGCTGTGCATGAGAGCGGGCGCCCAACGGGGAGAGCGGGTCCGT CCTCACGCAGCACAGCCCACCGAACACCGAGAGGAGAATATTTTTGC CGAAAACAGCCGCGGTCAACCGAGCAGTAGGATATTAGAATATGACGGTC GACACCCATTATCGTGTGCGACCCNATAAACACAACAACGATAGGCCNGA CAACTGATGAGACACAGCGCCGCGGAAGCGCAGCACGATGAGGCCGAAAG GAGACGACGACACGCACGCGCTACAGCGTGTCTACAACAAGTCGGCG GCGCGGCCAGAGCNTATTAGGCCTGGAGGATCGACCGCGAGATCTCACAA TGACTCACACATCACTCTACTTNTANGTATAGATAAAACAGAACAAAAAC GCAAAGTAATACTACAATGCGGAGCGTTACAGGATGGACGGCAGGCGATA CAGATGGACAGGACAGATCTGGAAAGGCAGTGACAGGTGTATAAGAGTAG ACAGTGCAGCGAAGAGTGAGGTCGACGAACAAGAAGAAGAAGATGCGAAAGA GTAGATGGAGATGAGTCGGAAACGAGAACATATGCGCGGNCGGACGGACT GAAAGAGAAATTGAGACAGTACAGGGTAGACCGAAGGTAGGCTGATAG ACCAGCAACACAGCACGCAGGAAGCGAATGAAGCACGCCGACGCGACGAG ACGAGCATAGAGCACACGCACACGACTGAGAGATCAGCACACAATGCGAA GGGCAGACCGCGGACGGCGCAGAAGCCGCCACCACACGTCCGGGACACCA CACGGGCACGGAGAACGAGGGAACACACGCGAGGAGGTACGATGAGGTGT GTGGACACGTGTCTGCTGGACGTAAAGAAGGTCAGGCGTAGCACGACACA GCAAACGGTGGGGCGCCCTAAACCGGTAGCGCCGAGAGACACGAAAAAG GACGGGCGCGCACGAGTTAGACGCAGACGATAGACAACCGGGACACAACA AAGGGGCAGATCGGAAGCACACGCGTGCACTAAGACAAGATAGGGCAATC CGATGCGGGAACGGCAGCGGACCGCAGTGCGAGACACAAAAAGCGGAGG GGAGAATAGCACCCCGACGCAATGAAAGAAGGGGTGTGAAAGAACCGCAT AAGGGTACCAATAACAGCATAGGAAAGAGTAGAGTACAGGAACAGCCAGA TGGCTGATGCAATATATGACCAAGTGGTGTCGTAGGCGCGGCGGTGAGCA GACCGGGGCGGAGGGCCGCGACCACAAGCGAGCGA

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>SARS120.CB_D10_-_078.ab1 1995 0 AGCGCGCCACGACACAGCGCGNATACGCACGCGACGACGCGGCGCGCGTCG CGCGCGCGGCGGTGGCGGCGCCGACGATCTCATCACGCTTCTGTTTGTC TNTCTTTGGCGGGGNATCAGCAGTCNTCTCGCTCTGTTTTTTTANNTTGG ATAATGCGCGCCCCCGTCGATGCGTCTGTNGTTATGCGCTTATTTATTA TCTCTGANAANNAACNAGGATGAGATGCATGCGGGCAGAGCTCTNATATA GGGACTCGTGGTGCACTCATGATCGTCGTACATCGAGCGACGTTTCCTCT ACCTCAATNGCAGCAGCCTAGCTTACTGGCTCGCGATACGATGCGCCGAC ATCATAACTCACGTANCCAATGTTCAGGCAAAGAGACGACGAGAGCGTGT AGTGATATCTCATGGCGATACTATACTGAGCAACGCGATGCATNGCTCGA TGGCGCGCGCGCACGCTCTAGCGACGCAGAATACTGCGAACGNAGAN CTACGAAATATGTACGATCTGTACATTGCTGCTGGTCGACTAATAACAGA TAAAGTCGACGTGTGTAGANACTATACTATTTAGCGGAGACCGAGTNC ACGTANTATACATGCGCATCGCTGACGCTGACGTCATATGATGNGCGACN GACTGCACGCACGCGCGCGCGTCNACTGCTGCGCACNGTCACTCGGGTG CCTGAGACGGTGATAGTAGTATATNCTATCACATTCACAGTGTCTATATT GATTATCATACGAGCATACGACGTGCAATTGCTGAGTGTCAGAGATGCGG GCGACGCCCACTACTGTCAGACGAGCAGTCATCATGCTGACGCCTGGTA CTTGCGAGACGACTGTATACNANTCGTGACGACATGCATGTATGACGNAT GACTTCAGTATATACGTCTGATTAGTGTGTCGCTANCGTACTTGCTGA CCTCAGTGAGCATACTACCACGATAGTACACTGGTATACACACTGATATA CGTAATACGATGAATGAAAGAAAGCAGACAGATGGTCGAGATGACGATAG TCATGTGCGTATTAGTCATGCACTACACGAGGTCCGGTCGCACTCAAGAT CCGTCCCTATANTGACAGCACGCACGACCACGCCTCACNACTCATATTTG TGATATCATATGTGAATTCTGACTGTCTCGTGCGTGATGACTACTANGTT GACTCGTCTCNGATGTACGTGCGGCCTGTGCGNACCGTGCGACTGTGCTA GTGTCANACGGCGCAACACATCTCACTGCGCGCATATCTGCTCGTCGTAC TGCGCANCATGCGCGACGTAGTCTAGACTTGGAGGNGAGTANCATATTGN GATCTGTGCGCAACACGTCGTGTGATCGTGACTCTNTCATGTTGTTGCT GCTCGTATNATCGATCNTCTNCACACTGTGTCGACGCATGATCACTATAG ATATGAGTAGCAGTGTCTGTTGGGTANACGANCACATGCGATACACGACC

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